10/576358

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<110> Smith, Austin, Gerard
 Ying, Qi-Long
 Nichols, Jennifer

<120> Improved Control Of ES Cell Self Renewal And Lineage Specification, And Medium Therefor

<130> 09641.0011-00000

<140> Not Yet Assigned

<141> 2006-04-17

<150> GB 0324270.8

<151> 2003-10-16

<150> GB 0324378.9

<151> 2003-10-17

<150> GB 0325007.3

<151> 2003-10-27

<160> 9

<170> PatentIn version 3.1

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<212> PRT

<213> Mus sp.

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Met Lys Ala Leu Ser Pro Val Arg Gly Cys Tyr Glu Ala Val Cys Cys 1 10 15

Leu Ser Glu Arg Ser Leu Ala Ile Ala Arg Gly Arg Gly Lys Ser Pro 20 25 30

Ser Thr Glu Glu Pro Leu Ser Leu Leu Asp Asp Met Asn His Cys Tyr 35 40 45

Ser Arg Leu Arg Glu Leu Val Pro Gly Val Pro Arg Gly Thr Gln Leu 50 55 60

Ser Gln Val Glu Ile Leu Gln Arg Val Ile Asp Tyr Ile Leu Asp Leu 65 70 75 80

Gln Val Val Leu Ala Glu Pro Ala Pro Gly Pro Pro Asp Gly Pro His 85 90 95

Leu Pro Ile Gln Thr Ala Glu Leu Thr Pro Glu Leu Val Ile Ser Lys
100 105 110

Asp Lys Arg Ser Phe Cys His 115

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<211> 119

<212> PRT

<213> Rattus sp.

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Met Lys Ala Leu Ser Pro Val Arg Gly Cys Tyr Glu Ala Val Cys Cys

1 10 15

Leu Ser Glu Arg Ser Leu Ala Ile Ala Arg Gly Arg Gly Lys Ser Pro 20 25 30

Ser Ala Glu Glu Pro Leu Ser Leu Leu Asp Asp Met Asn His Cys Tyr 35 40 45

Ser Arg Leu Arg Glu Leu Val Pro Gly Val Pro Arg Gly Thr Gln Leu 50 55 60

Ser Gln Val Glu Ile Leu Gln Arg Val Ile Asp Tyr Ile Leu Asp Leu 65 70 75 80

Gln Val Val Leu Ala Glu Pro Ala Pro Gly Pro Pro Asp Gly Pro His 85 90 95

Leu Pro Ile Gln Thr Ala Glu Leu Thr Pro Glu Leu Val Ile Ser Lys
100 105 110

Asp Lys Arg Ser Phe Cys His 115 <210> 3

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<213> Canis sp.

<400> 3

Met Lys Ala Leu Ser Pro Val Arg Gly Cys Tyr Glu Ala Val Cys Cys

1 10 15

Leu Ser Glu Arg Ser Leu Ala Ile Ala Arg Gly Arg Gly Lys Gly Pro
20 25 30

Ala Ala Glu Glu Pro Leu Ser Leu Leu Asp Asp Met Asn His Cys Tyr 35 40 45

Ser Arg Leu Arg Glu Leu Val Pro Gly Val Pro Arg Gly Thr Gln Leu 50 55 60

Ser Gln Val Glu Ile Leu Gln Arg Val Ile Asp Tyr Ile Leu Asp Leu 65 70 75 80

Gln Val Val Leu Ala Glu Pro Ala Pro Gly Pro Pro Asp Gly Pro His
85 90 95

Leu Pro Ile Gln Thr Ala Glu Leu Ala Pro Glu Leu Val Ile Ser Asn 100 105 110

Asp Lys Arg Ser Phe Cys His 115

<210> 4

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<213> Homo sapiens

<400> 4

Met Lys Ala Leu Ser Pro Val Arg Gly Cys Tyr Glu Ala Val Cys Cys
1 10 15

Leu Ser Glu Arg Ser Leu Ala Ile Ala Arg Gly Arg Gly Lys Gly Pro 20 25 30 Ala Ala Glu Glu Pro Leu Ser Leu Leu Asp Asp Met Asn His Cys Tyr 35 40 45

Ser Arg Leu Arg Glu Leu Val Pro Gly Val Pro Arg Gly Thr Gln Leu 50 55 60

Ser Gln Val Glu Ile Leu Gln Arg Val Ile Asp Tyr Ile Leu Asp Leu 65 70 75 80

Gln Val Val Leu Ala Glu Pro Ala Pro Gly Pro Pro Asp Gly Pro His 85 90 95

Leu Pro Ile Gln Thr Ala Glu Leu Ala Pro Glu Leu Val Ile Ser Asn 100 105 110

Asp Lys Arg Ser Phe Cys His

<210> 5

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<212> PRT

<213> Human immunodeficiency virus

<400> 5

Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg 1

<210> 6

<211> 16

<212> PRT

<213> Antennapedia

<400> 6

Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys 1 5 10 15

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<212> PRT

<213> Artificial Sequence

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<223> synthetic

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Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg Met Lys Ala Leu Ser 1 5 10 15

Pro Val Arg Gly Cys Tyr Glu Ala Val Cys Cys Leu Ser Glu Arg Ser 20 25 30

Leu Ala Ile Ala Arg Gly Arg Gly Lys Gly Pro Ala Ala Glu Glu Pro
35 40 45

Leu Ser Leu Leu Asp Asp Met Asn His Cys Tyr Ser Arg Leu Arg Glu 50 55 60

Leu Val Pro Gly Val Pro Arg Gly Thr Gln Leu Ser Gln Val Glu Ile 65 70 75 80

Leu Gln Arg Val Ile Asp Tyr Ile Leu Asp Leu Gln Val Val Leu Ala 85 90 95

Glu Pro Ala Pro Gly Pro Pro Asp Gly Pro His Leu Pro Ile Gln Thr 100 105 110

Ala Glu Leu Ala Pro Glu Leu Val Ile Ser Asn Asp Lys Arg Ser Phe 115 120 125

Cys His

<210> 8

<211> 135

<212> PRT

<213> Artificial Sequence

<220>

<223> synthetic

<400> 8

Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys 1 5 10 15

Met Lys Ala Leu Ser Pro Val Arg Gly Cys Tyr Glu Ala Val Cys Cys

20 25 30

Leu Ser Glu Arg Ser Leu Ala Ile Ala Arg Gly Arg Gly Lys Gly Pro 35 40 45

Ala Ala Glu Glu Pro Leu Ser Leu Leu Asp Asp Met Asn His Cys Tyr 50 55 60

Ser Arg Leu Arg Glu Leu Val Pro Gly Val Pro Arg Gly Thr Gln Leu 65 70 75 80

Ser Gln Val Glu Ile Leu Gln Arg Val Ile Asp Tyr Ile Leu Asp Leu 85 90 95

Gln Val Val Leu Ala Glu Pro Ala Pro Gly Pro Pro Asp Gly Pro His 100 105 110

Leu Pro Ile Gln Thr Ala Glu Leu Ala Pro Glu Leu Val Ile Ser Asn 115 120 125

Asp Lys Arg Ser Phe Cys His 130 135

<210> 9

<211> 135

<212> PRT

<213> Artificial sequence

<220>

<223> synthetic

<400> 9

Met Lys Ala Leu Ser Pro Val Arg Gly Cys Tyr Glu Ala Val Cys Cys 1 5 10 15

Leu Ser Glu Arg Ser Leu Ala Ile Ala Arg Gly Arg Gly Lys Ser Pro 20 25 30

Ser Thr Glu Glu Pro Leu Ser Leu Leu Asp Asp Met Asn His Cys Tyr 35 40 45

Ser Arg Leu Arg Glu Leu Val Pro Gly Val Pro Arg Gly Thr Gln Leu 50 55 60

Ser Gln Val Glu Ile Leu Gln Arg Val Ile Asp Tyr Ile Leu Asp Leu 65 70 75 80

Gln Val Val Leu Ala Glu Pro Ala Pro Gly Pro Pro Asp Gly Pro His
85 90 95

Leu Pro Ile Gln Thr Ala Glu Leu Thr Pro Glu Leu Val Ile Ser Lys
100 105 110

Asp Lys Arg Ser Phe Cys His Arg Gln Ile Lys Ile Trp Phe Gln Asn 115 120 125

Arg Arg Met Lys Trp Lys Lys 130 135